

Coordinate Regulation of TLR-Mediated Arachidonic Acid Mobilization in Macrophages by Group IVA and Group V Phospholipase A₂s¹

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Macrophages can be activated through TLRs for a variety of innate immune responses. In contrast with the wealth of data existing on TLR-dependent gene expression and resultant cytokine production, very little is known on the mechanisms governing TLR-mediated arachidonic acid (AA) mobilization and subsequent eicosanoid production. We have previously reported the involvement of both cytosolic group IVA phospholipase A₂ (cPLA₂) and secreted group V phospholipase A₂ (sPLA₂-V) in regulating the AA mobilization response of macrophages exposed to bacterial LPS, a TLR4 agonist. In the present study, we have used multiple TLR agonists to define the role of various PLA₂s in macrophage AA release via TLRs. Activation of P388D₁ and RAW2647.1 macrophage-like cells via TLR1/2, TLR2, TLR3, TLR4, TLR6/2, and TLR7, but not TLR5 or TLR9, resulted in AA mobilization that appears to involve the activation of both cPLA₂ and sPLA₂ but not of calcium-independent phospholipase A₂. Furthermore, inhibition of sPLA₂-V by RNA interference or by two cell-permeable compounds, namely scalaradial and manoalide, resulted in a marked reduction of the phosphorylation of ERK1/2 and cPLA₂ via TLR1/2, TLR2, TLR3, and TLR4, leading to attenuated AA mobilization. Collectively, the results suggest a model whereby sPLA₂-V contributes to the macrophage AA mobilization response via various TLRs by amplifying cPLA₂ activation through the ERK1/2 phosphorylation cascade. *The Journal of Immunology*, 2009, 182: 3877–3883.

Phospholipase A₂ (PLA₂)³ enzymes hydrolyze membrane phospholipids at the sn-2 position of the glycerol backbone, releasing a free fatty acid and a lysophospholipid. This reaction is particularly important when the fatty acid liberated is arachidonic acid (AA), which can be converted into biologically active compounds called the eicosanoids (1–5). In the latest update, the PLA₂ enzymes were classified into 15 group types, according to their primary sequence (1). Additionally, a 16th PLA₂ group has been reported very recently (6). However, a second classification of the PLA₂ enzymes, sometimes more useful, also exists that categorizes the enzymes into five major families attending to biochemical commonalities (2, 7). These families are the Ca²⁺-dependent secreted enzymes, the Ca²⁺-dependent cytosolic enzymes, the Ca²⁺-independent cytosolic enzymes, the platelet-activating factor acetyl hydrolases, and the lysosomal PLA₂s. Of these families, the first two have been repeat-

edly implicated in AA mobilization in response to a variety of immunoinflammatory stimuli (1–5). Today, there is general consensus that the calcium-dependent cytosolic group IVA PLA₂ (cPLA₂) is the critical enzyme in AA release (8, 9) and that, depending on cell type and stimulation conditions, a secreted PLA₂, in particular that belonging to groups IIA, V and X, may also participate by creating an amplification loop (10–14).

TLRs are a family of structural proteins in mammals related to Toll receptors in *Drosophila*. TLRs play a central role in inflammation and innate immune defense by recognizing specific structures of pathogens (15, 16). To date, 13 members of the TLR family have been identified in mice. TLR4, the first mammalian TLR identified and the best characterized member of this family (15–18), has been established as an essential component in the recognition of Gram-negative bacterial LPS. Additionally, TLR4 recognizes several other ligands. TLR2 recognizes a variety of microbial components and cooperates with TLR1 or TLR6 to recognize triacyl and diacyl lipopeptides, respectively. TLR5 recognizes bacterial flagellin. TLR3 recognizes dsRNA, and TLR7 and TLR9 recognize single-stranded nucleic acids. (15–18).

In previous work we have extensively investigated the mechanism of PLA₂-mediated AA mobilization in P388D₁ macrophage-like cells stimulated via TLR4 by bacterial LPS (19–26). In the present work, we have extended these studies to other TLRs. Our studies highlight the differential contributions of cPLA₂ and secreted group V phospholipase A₂ (sPLA₂-V) to the AA mobilization response and, in addition, suggest a role for sPLA₂-V in regulating ERK-dependent cPLA₂ phosphorylation in response to stimulation of some, but not all, TLRs.

Materials and Methods

Reagents

P388D₁ macrophage-like cells (MAB clone) (19) were provided by Dr. Ed Dennis (University of California at San Diego, La Jolla, CA). RAW 264.7 macrophage-like cells were obtained from the American Type Culture Collection. RPMI 1640 medium and FBS were from Invitrogen.

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Received for publication December 2, 2008. Accepted for publication January 15, 2009.

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¹ This work was supported by the Spanish Ministry of Science and Innovation (Grants SAF2007-60055 and BFU2007-67154).

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³ Abbreviations used in this paper: PLA₂, phospholipase A₂; AA, arachidonic acid; cPLA₂, group IVA cytosolic phospholipase A₂; sPLA₂, secreted phospholipase A₂; sPLA₂-V, group V secreted phospholipase A₂; BEL, bromoenol lactone; Pam3SCK4, N-palmitoyl-5-[2,3-bis(palmitoyloxy)-(2R,2S)-propyl]-Cys-[S]-Ser-[S]-Lys(4) trihydrochloride; LTA, lipoteichoic acid from *Staphylococcus aureus*; FSL-1, S-(2,3-bis-palmitoyloxypropyl)-Cys-Gly-Asp-Pro-Lys-His-Pro-Ser-Phe; ODN1826, synthetic oligodeoxynucleotide 1826, 5'-TCCATGACGTTCTGACGTT-3'; siRNA, small interfering RNA; iPLA₂, calcium-independent phospholipase A₂.

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[5,6,8,9,11,12,14,15-³H]AA (sp. act. 214 Ci/mmol) was from GE Healthcare. LPS (*Escherichia coli* 0111:B4), poly(I:C), and bromoanil lactone (BEL), were from Sigma-Aldrich. 12-Epi-scalarial and manoolide were from BIOMOL. The TLR agonists *N*-palmitoyl-S-[2,3-bis(palmitoyloxy)-(2*R*,2*S*)-propyl]-Cys-[S]-Ser-[S]-Lys(4) trihydrochloride (Pam3CSK4), lipoteichoic acid from *Staphylococcus aureus* (LTA), flagellin from *Salmonella typhimurium*, S-(2,3-bispalmitoyloxypropyl)-Cys-Gly-Asp-Pro-Lys-His-Pro-Ser-Phe (FSL-1), imiquimod, and synthetic oligodeoxynucleotide 1826 (5'-TCCATGACGTTCTGACGTT-3'; ODN1826) were purchased from InvivoGen. Rabbit polyclonal Abs that detect activating phosphorylations of ERK1/2 (Thr²⁰²/Tyr²⁰⁴) and cPLA₂ (Ser⁵⁰⁵) were from Cell Signaling Technology. All other reagents were from Sigma-Aldrich.

Cell culture and labeling conditions

P388D₁ and RAW264.7 cells were maintained at 37°C in a humidified atmosphere at 95% air and 5% CO₂ in RPMI 1640 medium supplemented with 5% and 10% FBS, respectively, 2 mM glutamine, 100 U/ml penicillin, and 100 µg/ml streptomycin. Cells were plated at 10⁶/well, allowed to adhere overnight, and used for experiments the following day. All experiments were conducted in serum-free RPMI 1640 medium. When required, radiolabeling of the cells was achieved by including 0.25 µCi/ml during the overnight adherence period (20 h). Labeled fatty acid that had not been incorporated into cellular lipids was removed by washing the cells twice with serum-free medium containing 0.5 mg/ml albumin. Stimuli were added for 20 h at the following doses: LPS, 100 ng/ml; poly(I:C), 25 µg/ml; Pam3CSK4, 1 µg/ml; LTA, 1 µg/ml; flagellin, 50 ng/ml; FSL-1, 1 µg/ml; imiquimod, 5 µg/ml; ODN1826, 1 µM. Preliminary experiments indicated that at the above-specified concentrations, the inhibitors induced maximal responses. When inhibitors were used they were added 30 min before the stimuli. In phosphorylation assays, TLR agonists were added for 30 min.

Small interfering RNA (siRNA) inhibition studies

The siRNA directed against sPLA₂-V was from Eurofins MWG Operon (sequence 5'-CAC GAC UCC UUC UGU CCA AdTdT-3'). The cells (3 × 10⁵/ml) were transiently transfected with oligonucleotide (5–20 nM) in the presence of 10 µg/ml Lipofectamine (Invitrogen) under serum-free conditions for 6 h. Afterward, 5% serum was added and the cells were maintained at normal culture conditions for 20 h. Afterward, the cells were used for experiments as described above. A scrambled siRNA was used as a negative control.

Immunoblot analyses

Cells were serum-starved and stimulated in the absence or presence of the indicated inhibitors. Afterward, the cells were washed and then lysed in a buffer consisting of 20 mM Tris-HCl (pH 7.4), 150 mM NaCl, 0.5% Triton X-100, 100 µM Na₂VO₄, 1 mM PMSF, 10 mg/ml aprotinin, and 10 mg/ml leupeptin at 4°C. Protein was quantified, and a 50-µg aliquot for p-ERK1/2 and p-cPLA₂ was separated by SDS/PAGE (10% acrylamide) and transferred to Immobilon-P (Millipore). Nonspecific binding was blocked with 5% nonfat in PBS and then incubated with specific Abs. Bands were detected by the Amersham ECL system.

Measurement of extracellular fatty acid release

The cells were placed in serum-free medium for 1 h before the addition of TLR agonists in the absence or presence of inhibitors and in the presence of 0.5 mg/ml BSA to blunt fatty acid reacylation (27). After the 24-h incubation period, the supernatants were removed, cleared of detached cells by centrifugation, and assayed for radioactivity by liquid scintillation counting.

PLA₂ activity measurements

PLA₂ activity assays were conducted as previously described (28–35). Briefly, for calcium-independent PLA₂ (iPLA₂) activity, homogenates from cells treated with 10 µM BEL for 30 min were prepared, and 100-µg aliquots were incubated for 2 h at 37°C in 100 mM HEPES (pH 7.5) containing 5 mM EDTA and 100 µM labeled phospholipid substrate, 1-palmitoyl-2-[³H]palmitoyl-*sn*-glycero-3-phosphocholine, in a final volume of 150 µl. The phospholipid substrate was used in the form of sonicated micelles in buffer. The reactions were quenched by adding 3.75 volumes of chloroform/methanol (1:2). After lipid extraction, free [³H]palmitic acid was separated by thin-layer chromatography using *n*-hexane/ethyl ether/acetic acid (70:30:1, v/v/v) as the mobile phase. For measuring sPLA₂ activity, the mammalian membrane assay described by Diez et al. (36) was used. Briefly, aliquots of U937 cell homogenates were incubated for 1–2 h at 37°C in 100 mM HEPES (pH 7.5) containing 1.3

Table I. TLR agonists used in this study

Agonist	Description	TLR
Pam3CSK4	<i>N</i> -palmitoyl-S-[2,3-bis(palmitoyloxy)-(2 <i>R</i> ,2 <i>S</i>)-propyl]-Cys-[S]-Ser-[S]-Lys(4)	1/2
LTA	Lipoteichoic acid from <i>Staphylococcus aureus</i>	2
Poly(I:C)	Polyinosinic-polycytidylic acid	3
LPS	LPS from <i>Escherichia coli</i> 0111:B4	4
Flagellin	Flagellin from <i>Salmonella typhimurium</i>	5
FSL-1	S-(2,3-bispalmitoyloxypropyl)-Cys-Gly-Asp-Pro-Lys-His-Pro-Ser-Phe	6/2
Imiquimod	1-isobutyl-1 <i>H</i> -imidazo[4,5- <i>c</i>]quinolin-4-amine	7
ODN1826	5'-TCCATGACGTTCTGACGTT-3'	9

mM CaCl₂ and 100,000 dpm of [³H]AA-labeled U937 cell membrane, used as substrate, in a final volume of 0.15 ml. Before assay, the cell membrane substrate was heated at 57°C for 5 min to inactivate CoA-independent transacylase activity. The assay contained 1 µM pyrrophenone and 25 µM BEL to completely inhibit endogenous cPLA₂ and iPLA₂ activities (37, 38). After lipid extraction, free [³H]AA was separated by thin-layer chromatography, using *n*-hexane/ethyl ether/acetic acid (70:30:1, v/v/v) as a mobile phase.

Other methods

Protein concentration was determined utilizing the Bradford protein assay kit (Bio-Rad) with BSA as a standard. Densitometric quantitation of bands was analyzed by Quantity One program (Bio-Rad).

Data presentation

Unless otherwise stated, data are presented as the means ± SEM of at least three different experiments.

Results

TLR-mediated AA mobilization in P388D₁ cells

We have previously reported that both cPLA₂ and sPLA₂-V participate in the AA mobilization response of P388D₁ macrophage-like cells to LPS, a stimulus that signals through TLR4 (19–26). In the present work we have extended our studies to other TLRs, of which very little information is available with regard to AA mobilization and eicosanoid production. Thus, we began this study by analyzing the AA release response of P388D₁ macrophage-like cells to a variety of TLR agonists, which are listed in Table I. In keeping with our previous data (19–26), engagement of TLR4 receptors by LPS induced abundant AA mobilization. Additionally, stimulation via TLR1/2 (Pam3CSK₄), TLR2 (LTA), TLR3 (poly(I:C)), TLR6/2 (FSL-1), and TLR7 (imiquimod) also significantly enhanced AA release (Fig. 1A). However, stimulation of cells via TLR5 (flagellin) or TLR9 (ODN1826) had little or no effect on the response (Fig. 1A). To ensure that these results are physiologically relevant, we sought to discard unwanted cell-specific effects by conducting the same measurements on another murine macrophage-like cell line from a different genetic background. We used the cell line RAW264.7, which is derived from BALB/c mice. The P388D₁ cell line is derived from DBA/2 mice. Both cell lines are mature murine macrophage cell lines with similar phenotypic and functional characteristics (39–41). Furthermore, we have characterized by PCR the expression profile of PLA₂ isoforms potentially involved in AA release in these cell lines and found it to be comparable, that is, both cell lines express group V, group IVA, and group VIA PLA₂s, but not group IIA or group X PLA₂s (V. Ruipérez, Y. Sáez, and J. Balsinde, unpublished results). Compared with P388D₁ cells, activated RAW264.7 cells

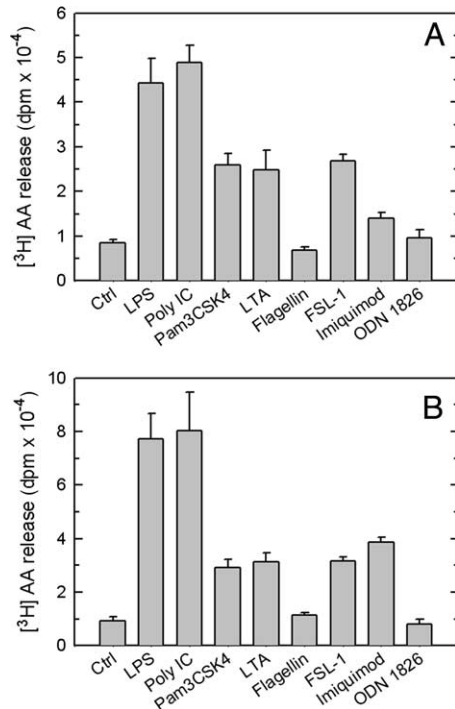


FIGURE 1. TLR-mediated AA mobilization in macrophage-like cell lines. P388D₁ (A) or RAW 264.7 (B) cells, labeled with [³H]AA, were incubated with various TLR agonists for 20 h at the concentrations indicated in *Materials and Methods*. [³H]AA in the extracellular media was estimated by scintillation counting.

generally released higher amounts of AA to the extracellular medium. However, qualitatively, the results were very similar in both cell lines, that is, significant AA release in response to agonists of TLR1/2, TLR2, TLR3, TLR4, TLR6/2, and TLR7, and no effect to agonists of TLR5 and TLR9 (Fig. 1B).

PLA₂ forms potentially capable of catalyzing AA mobilization are usually categorized into three major families, namely the Ca²⁺-dependent cPLA₂s, the sPLA₂s, and the iPLA₂s (2, 7). Macrophage cells are known to express all three classes of PLA₂ (37). Thus, a useful way to initially distinguish among the PLA₂s is the use of chemical inhibitors that selectively inhibit each of the forms. To this end, we utilized pyrrophenone (cPLA₂ inhibitor), scalaradial (sPLA₂ inhibitor), and BEL (iPLA₂ inhibitor) (38). In previous work we have extensively characterized the effect of these inhibitors on cells and demonstrated that, at the concentrations employed in this work, none of the inhibitors interferes with PLA₂ activities distinct from those against which the inhibitors are directed (26, 28–35). Table II shows that 1 μM pyrrophenone almost

Table II. Effect of PLA₂ inhibitors on TLR-mediated AA release in P388D₁ cells^a

Agonist	Pyrrophenone	Scalaradial	BEL
Pam3CSK4	1 ± 1	52 ± 1	100 ± 6
LTA	2 ± 1	49 ± 3	100 ± 4
Poly(I:C)	6 ± 1	76 ± 1	89 ± 4
LPS	7 ± 2	77 ± 2	92 ± 6
FSL-1	2 ± 1	50 ± 1	85 ± 4
Imiquimod	5 ± 1	62 ± 3	100 ± 1

^a The cells, labeled with [³H]AA, were treated with 1 μM pyrrophenone, 4 μM scalaradial, or 10 μM BEL, as indicated, for 30 min. Afterward the cells were incubated with the indicated TLR agonists for 20 h. AA in the supernatants was measured by scintillation counting. The results are expressed as a percentage of the response observed to each stimulus in the absence of inhibitor.

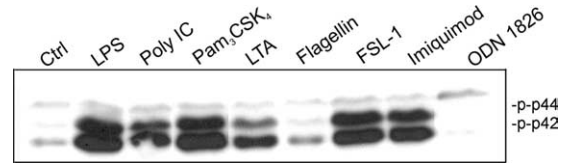


FIGURE 2. ERK1/2 phosphorylation in response to TLR agonists. P388D₁ cells were treated with TLR agonists for 30 min at the concentrations indicated in *Materials and Methods*, and ERK1/2 phosphorylation was determined by immunoblot.

completely ablated the AA release response to all TLR agonists used, pointing to cPLA₂ as the key regulatory enzyme. Interestingly, the sPLA₂ inhibitor scalaradial also produced significant inhibitory responses, albeit much lower than those elicited by pyrrophenone (Table II). On the other hand, the iPLA₂ inhibitor BEL exerted no significant effects on any of the agonists tested (Table II), suggesting no role for iPLA₂ in TLR-mediated AA mobilization. Since the BEL experiments produced negative results, and to assess whether the inhibitor was indeed acting on the iPLA₂ under these conditions, we measured as a control the iPLA₂ activity of homogenates from BEL-treated cells. We found a 70 ± 6% reduction of endogenous iPLA₂ activity in BEL-treated homogenates with respect to that measured in homogenates from untreated cells. Thus, our inability to detect any effect of iPLA₂ inhibition on TLR-mediated AA release under conditions where cellular iPLA₂ activity is effectively blunted demonstrates the lack of involvement of iPLA₂ in the response under study.

sPLA₂ modulates ERK and cPLA₂ phosphorylation during stimulation via TLRs

In the next series of experiments, we directed our attention toward the enzymes that activate cPLA₂. Phosphorylation of cPLA₂ on Ser⁵⁰⁵ results in an ~2–2.5-fold increase in its specific activity (42). Many of the data currently available in the literature suggest that cPLA₂ is phosphorylated and activated by either the ERK1/2 (p44/p42) or p38 isoforms of MAPK; the activation mechanism may vary with the cell type and activation regimen (42). In this regard, we have previously shown that phosphorylation activation of the AA-releasing cPLA₂ at Ser⁵⁰⁵ occurs during stimulation of P388D₁ macrophage-like cells by various agonists, including the TLR4 agonist LPS, and that this phosphorylation is effected by the ERKs p42 (ERK1) and p44 (ERK2) (43, 44). Fig. 2 shows that these two kinases are activated in P388D₁ cells stimulated not only via TLR4, but also via TLR1/2, TLR2, TLR3, TLR6/2, and TLR7. In our hands, agonists of TLR5 and TLR9 did not have any effect. Note that this pattern of ERK phosphorylation activation by TLR agonists is identical to that observed previously for AA mobilization (Fig. 1). Interestingly, when the ERK phosphorylation measurements were conducted in cells incubated with 4 μM scalaradial to block sPLA₂ activity, a clear inhibition of the phosphorylation of the ERKs was detected in response to stimulation via TLR1/2, TLR2, TLR3, and TLR4 (Fig. 3A). Intriguingly, phosphorylation activation of the ERKs via TLR6/2 and TLR7 was not appreciably changed by the presence of scalaradial (Fig. 3A), which highlights a selectivity of action of the drug. Thus, depending on the agonist, there appear to be scalaradial-dependent and -independent pathways for ERK phosphorylation within the same cell type. More importantly, the phosphorylation of cPLA₂ was also found to be inhibited in scalaradial-treated cells stimulated via TLR1/2, TLR2, TLR3, and TLR4 (Fig. 3A).

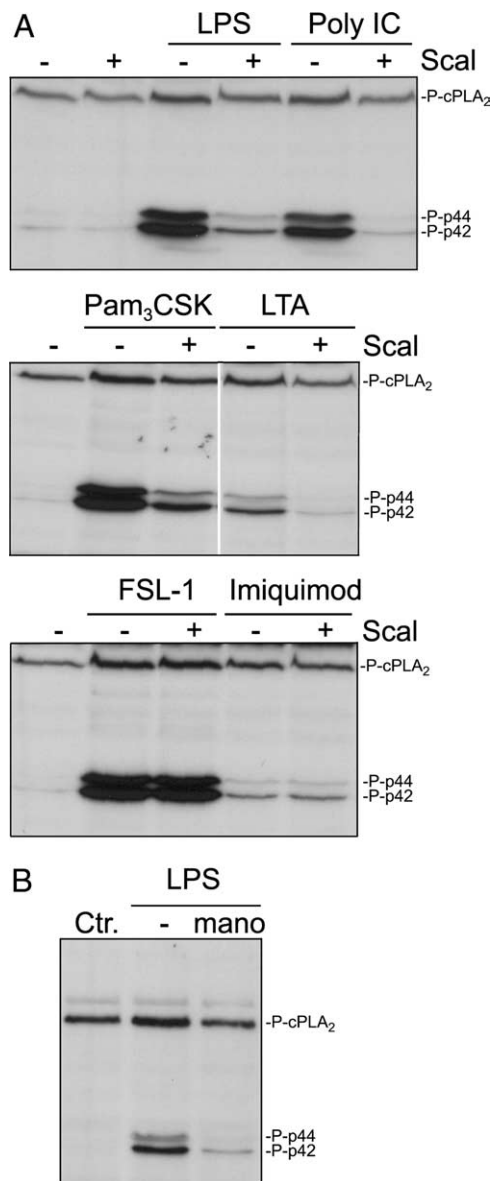


FIGURE 3. Effect of sPLA₂ inhibition on ERK1/2 and cPLA₂ phosphorylation. *A*, P388D₁ cells were treated with (+) or without (–) 4 μM scalaradial (Scal) for 30 min, as indicated. Afterward the cells were exposed to the TLR agonists for 30 min at the concentrations indicated in *Materials and Methods*, and the phosphorylation of ERK1/2 and cPLA₂ was determined by immunoblot. *B*, P388D₁ cells were treated with 4 μM manoalide, exposed to LPS for 30 min, and the phosphorylation of ERK1/2 and cPLA₂ was determined by immunoblot.

The effect of another sPLA₂ inhibitor, manoalide (26), on TLR-dependent phosphorylation of ERKs and cPLA₂ is shown in Fig. 3*B*. This compound, at concentrations that do not interfere with cellular cPLA₂ activity as judged by *in vitro* assay (26), also promoted a marked inhibition of the phosphorylation reactions. Because manoalide is structurally unrelated to scalaradial, these data provide additional independent evidence for the role of cellular sPLA₂ in regulating the phosphorylation state of cPLA₂.

Collectively, the results shown in Fig. 3, obtained with two structurally unrelated sPLA₂ inhibitors, are suggestive of sPLA₂ activity regulating ERK activation by TLRs and, in turn, cPLA₂ phosphorylation and enhanced AA release. However, studies using chemical inhibitors are limited in their ability to

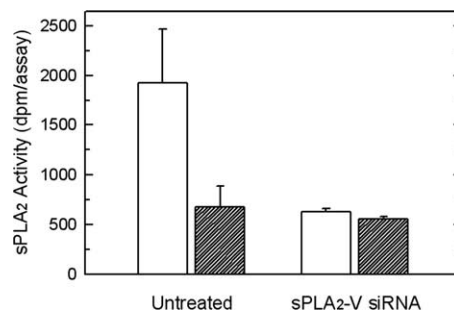


FIGURE 4. Effect of siRNA targeting sPLA₂-V on sPLA₂ activity. P388D₁ cells were treated or not with siRNA targeting sPLA₂-V, as indicated, and sPLA₂ activity was measured in cell homogenates from LPS-treated cells in the absence (open bars) or presence (dashed bars) of 4 μM scalaradial.

identify a particular group of sPLA₂. To confirm the molecular identity of the sPLA₂ and, in turn, to obtain more definite evidence for the role of sPLA₂ in regulating cPLA₂, we utilized siRNA technology to selectively knock down the expression of sPLA₂-V, the enzyme that accounts for most of the sPLA₂ activity of P388D₁ cells (45), and also the form that has been implicated by us and others as being involved in AA mobilization in cells relevant to innate immunity, such as murine macrophages and mast cells (46, 47). Since we have been unable to find reliable Abs against murine sPLA₂-V, the efficiency of siRNA knockdown was judged by enzyme activity assay in cell homogenates. As shown in Fig. 4, homogenates from cells deficient in sPLA₂-V showed a marked reduction in sPLA₂ activity as compared with negative controls. Importantly, when the effect of scalaradial was studied in homogenates from sPLA₂-V-deficient cells, no further inhibition of activity was detected, providing experimental support to the idea that sPLA₂-V contributes to most of the sPLA₂ activity of P388D₁ cells (44). In turn, the data of Fig. 4 provide additional evidence of the selectivity of action of scalaradial on sPLA₂-V in P388D₁ macrophage-like cells.

Blockade of sPLA₂-V expression by siRNA resulted in the inhibition of the phosphorylation of the ERKs in response to stimulation via TLR4 (Fig. 5), and also in the phosphorylation of cPLA₂ (Fig. 5). Collectively, these results suggest that, in addition to any direct effect of sPLA₂-V on cellular phospholipids,

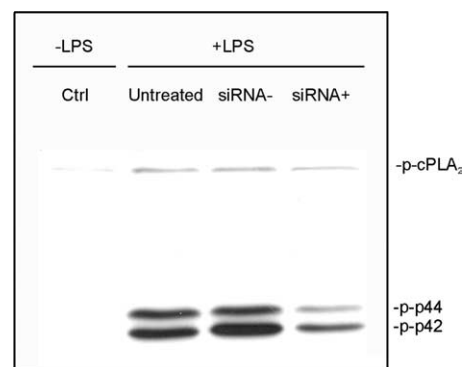


FIGURE 5. Inhibition of phosphorylation of ERK1/2 and cPLA₂ by siRNA targeting sPLA₂-V in TLR4-stimulated cells. Cells were either untreated or transfected with a siRNA negative control (siRNA⁻) or siRNA targeting sPLA₂-V (siRNA⁺). Afterward, the cells were incubated with or without 100 ng/ml LPS for 30 min, as indicated, and the phosphorylation of ERK1/2 and cPLA₂ was determined by immunoblot.

sPLA₂-V may participate in TLR-mediated AA mobilization by modulating the phosphorylation state of cPLA₂.

Discussion

TLRs are a class of receptors on the surface of immune cells that recognize structurally conserved molecules derived from pathogens (15–18). TLRs interact with different combinations of adapter proteins and activate transcription factors leading to specific immune responses (15–18). In contrast to the wealth of data that exist on TLR-dependent gene expression and resultant cytokine expression, very little is known on the mechanisms governing TLR-mediated AA mobilization and eicosanoid production. Most of what is known in this regard has been obtained from studies using the TLR4 agonist LPS as a model stimulus (48).

Aside from TLR4, there are very limited data on the AA mobilization mechanisms triggered by engagement of other TLRs. Here we have comparatively investigated the effectors involved in the AA release response triggered by various TLRs in murine macrophages. An unexpected finding is that not all TLRs couple to stimulation of AA mobilization. Stimulation of the cells via TLR5 and TLR9 fail to promote significant AA mobilization, an observation that was made utilizing two different cell lines, namely P388D₁ and RAW264.7. It seems likely that cellular stimulation of the cells by TLR5 or TLR9 fails to recruit an adapter/regulatory protein that is crucial for eliciting the AA release response. Studies are ongoing in our laboratory to identify regulatory components of TLR signaling that are not recruited after TLR5/TLR9 occupancy but may be involved in signaling via other TLRs.

Our inability to detect significant AA mobilization via TLR9 in two macrophage cell lines is at variance with a very recent study by Lee et al. (49), who reported significant AA release via TLR9 in RAW264.7 cells. At present we have no explanation for this discrepancy, and we speculate that it may be due to differences in the experimental conditions utilized. It has been demonstrated that IFN- γ can sensitize murine macrophages to produce various cytokines in response to TLR9 stimulation (50–52). This raises the intriguing possibility that a “priming step” could be necessary for the cells to mobilize AA in response to TLR9 agonists. Regarding the lack of effect of TLR5 agonists on AA mobilization in our hands, another recent report by Buczynski et al. (53) also failed to detect eicosanoid production via stimulation of this receptor by flagellin in RAW264.7 macrophage-like cells.

On the other hand, stimulation of TLR1/2, TLR2, TLR3, TLR4, TLR6/2, and TLR7 by the appropriate stimuli results in significant AA mobilization, although some stimuli behave as more potent than others. In particular, stimulation via TLR3 and TLR4 yields very robust responses. A pharmacological analysis of PLA₂ effectors potentially involved in the response clearly reflected the participation of cPLA₂ and sPLA₂ but not of iPLA₂. It is striking that the pharmacological profile was identical for all the TLRs, that is, inhibition of cPLA₂ by pyrrophenone almost ablates the AA responses to all agonists, inhibition of sPLA₂ by scalaradial partially blunts it, and inhibition of iPLA₂ by BEL has no significant effect. Our data are fully consistent with recent data in murine peritoneal macrophages and mast cells showing that disruption of the cPLA₂ gene nearly abrogates eicosanoid biosynthesis, while disruption of the sPLA₂-V gene leads to a 35–50% reduction of the response (54–56). Thus, our data are in line with a scenario whereby cPLA₂ acts as the main effector of AA release, and sPLA₂ acts to amplify the cPLA₂-mediated response (57, 58).

The involvement of two different kinds of PLA₂, namely sPLA₂ and cPLA₂, raises the question of whether there is coordinated action between the two enzymes in regulating AA release via engagement of TLR receptors. A major mechanism for the regulation of cPLA₂ is the phosphorylation of the enzyme at Ser⁵⁰⁵, which results in an increased ability of the enzyme to hydrolyze membrane phospholipids (8, 42). Depending on cell type and/or stimulus, various members of the MAPK family may catalyze cPLA₂ phosphorylation (8, 42). In the P388D₁ macrophage-like cell system we previously showed that phosphorylation of cPLA₂ at Ser⁵⁰⁵ occurs during stimulation of the cells via TLR4, and that this phosphorylation is mediated by the ERKs p42 (ERK1) and p44 (ERK2) (43). We demonstrate in this work that inhibition of sPLA₂-V expression and activity by either siRNA or chemical inhibitors results in reduced phosphorylation of the ERKs and cPLA₂ in response to some but not all TLRs. Thus, our results suggest that sPLA₂-V participates in the AA mobilization response of various TLRs, namely TLR1/2, TLR2, TLR3, and TLR4, by modulating the phosphorylation state of cPLA₂ and, hence, its enzymatic activity. This is a relevant observation, since cross-talk between sPLA₂ and cPLA₂ enzymes in eliciting AA release has been repeatedly suggested in many instances, but a clear picture of this interaction is yet to emerge (5).

There is ample evidence that exogenous sPLA₂s and/or sPLA₂s overexpressed in various cell types are capable of amplifying the essential role of cPLA₂ in AA mobilization and subsequent eicosanoid synthesis (11, 14, 25, 59–61). sPLA₂-V could potentially be involved in cPLA₂-dependent AA mobilization through three pathways, one involving reinternalization via caveolin-rich domains (24, 62), the second involving direct interaction with phosphatidylcholine-rich outer membrane domains (62–64), and the third one involving an undefined intracellular action prior to secretion of the enzyme (65, 66). On the other hand, very limited evidence is available on the role of “endogenous”, preexisting sPLA₂. In this regard, our results provide a molecular mechanism whereby sPLA₂-V modulates the activity of cPLA₂ by regulating its phosphorylation via ERK, which is in agreement with recent data in murine mast cells (58). Thus, in addition to any direct effect on phospholipid hydrolysis per se, endogenous sPLA₂-V may participate in the AA mobilization responses mediated by some TLRs by regulating the sequential phosphorylation of ERK1/2 and cPLA₂.

Acknowledgments

We thank Montse Duque and Yolanda Sáez for expert technical assistance. Centro de Investigación Biomédica en Red de Diabetes y Enfermedades Metabólicas Asociadas is an initiative of the Instituto de Salud Carlos III.

Disclosures

The authors have no financial conflicts of interest.

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